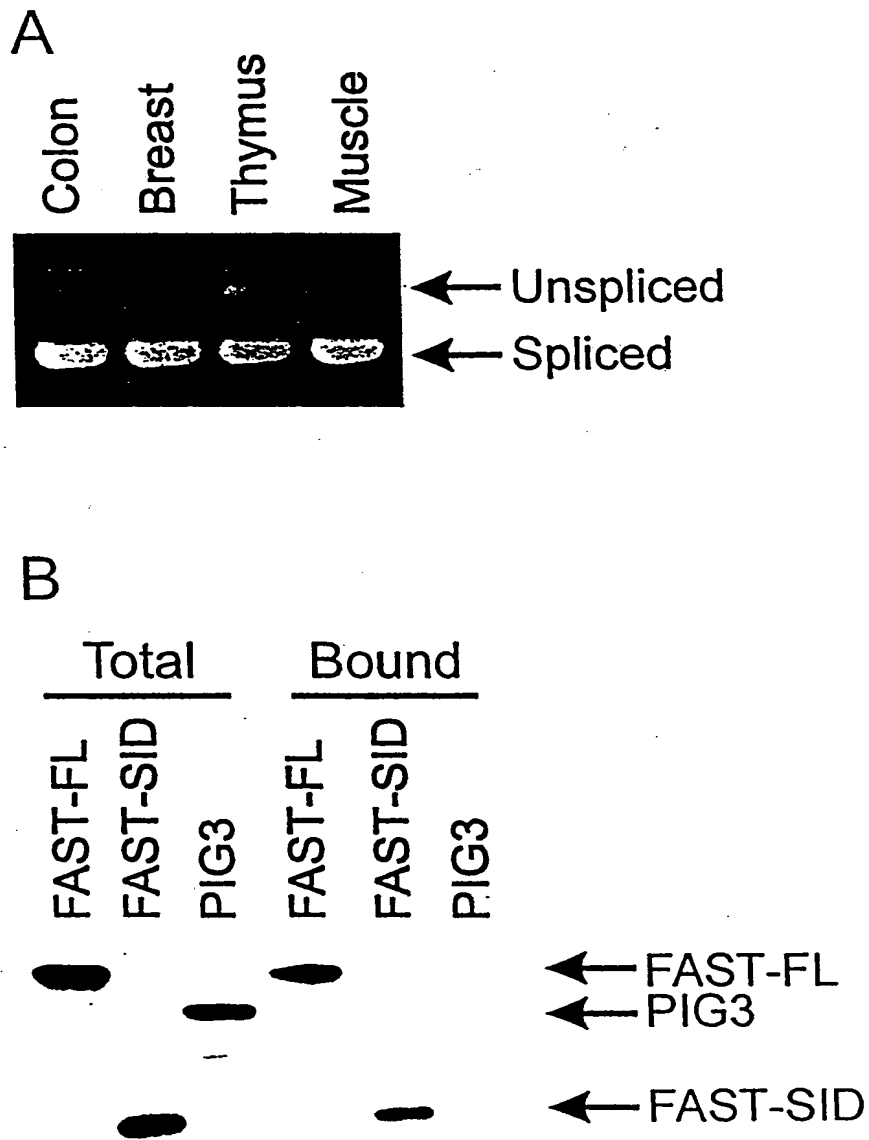


FIG. 1

ST-1	mgpcsgsrlgppaeaspsqpp-----	21
ST-1	mrdpslslyseffagsgyesveppslallssidqeqlpvatgqsynhsvqpw	51
ST-1	-----	21
ST-1	pqpwpplslyreggtwspdrsgmyglspgthegscththegpkdsmagdht	102
ST-1	---RRRRLRLHDPPYTYLMTLCAAFSRRLAOLIROQAVEPE	69
ST-1	rsrSKKENYHYPPYSYLMELVONSPEKRLSOTLKESTLEPE	153
ST-1	FREDYEGWIDSERHNLSSNRCFRYPKDEAPQAGNFWADYSLPEAEAL	120
ST-1	ENGDMGWKDSERHNLSSSDCEKLLADEGAPQAGNETTIDYSLPELDEM	204
ST-1	RKONTAECRwqngCARGAFAKDEGPYELHGRPYrppspppppsegffiks	171
ST-1	KKONTAEKTR-----GGSDYEVQDLAPYELHGRPYrppspppppsegffiks	250
ST-1	lggsgegagwpglapqsspvpagtgnsggeaaptpplpsserplwplcpl	222
ST-1	slsdaedsqqnttgkklntsfmidslldlqetldodasrnlengrispav	301
ST-1	pgptrvegetvqggaigpstlspeprawlqllqgtavpggrssgghrasl	273
ST-1	amnmwssaplllythsketrnarseglstinstyssssssistispvgfqk	352
ST-1	wgq-----	276
ST-1	eqeksgqrqtqrvghpikrsredddcsttssdpdtgnyspieppkkmplls	403
ST-1	-PPPSYLPIYTPNEMElappotscpcpstspaywgvapetrpgpgllc-	325
ST-1	dlPPPSYTKSVAENMELAPpsvleffhfprftyynygpspymtppywgfphpt	454
ST-1	-----DLDALFQGVPPNLSMDVWVSHPRDLAABpgpgwllsw	362
ST-1	nsggdsprgpqspLDLDNMLRAMPPNLSMDVWVSHPRDLAABpgpgwllsw	505
ST-1	cs1-----	365
ST-1	ssgspypsrqglm	518

FIG. 2



A

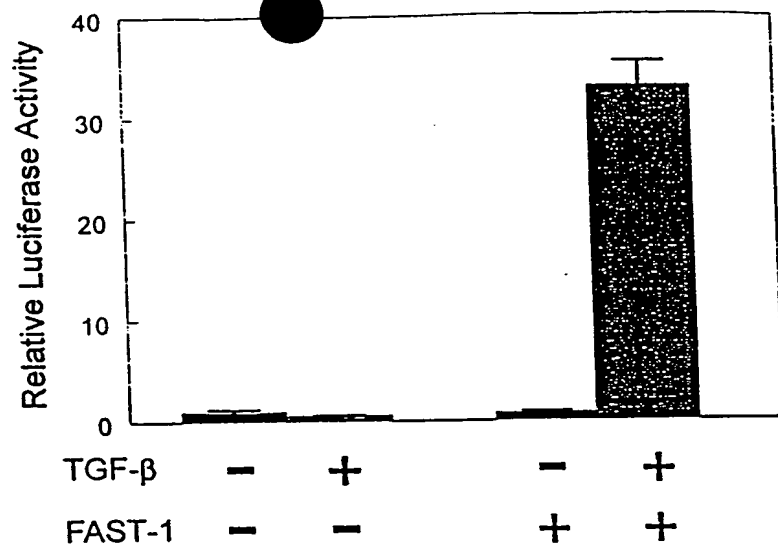
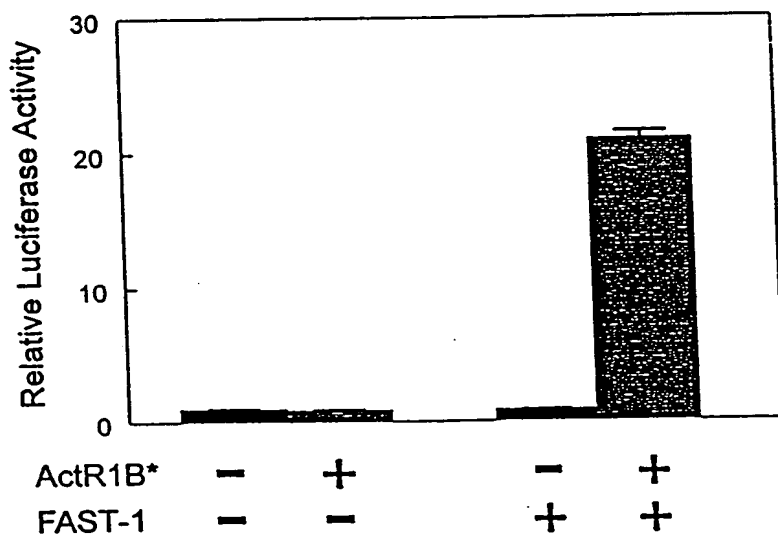


FIG. 3

B



C

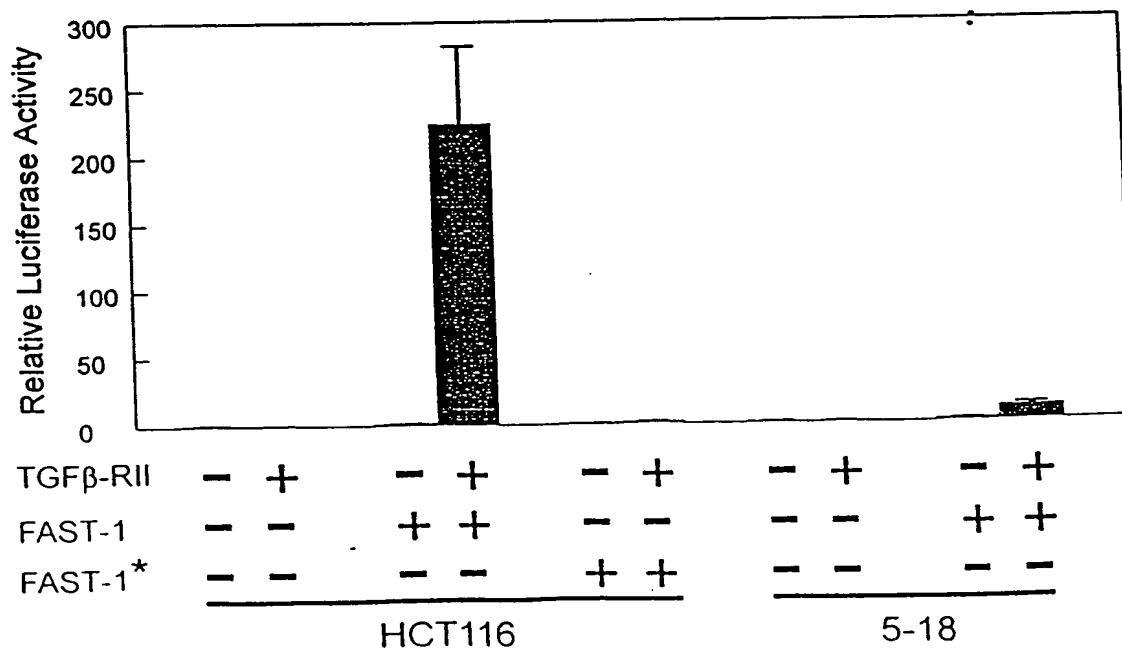


FIG. 4



CONSENSUS									
	T	G	T	GT	GT	A	T	T	
G	41	41	11	0	54	47	0	0	70
A	23	35	17	0	5	0	0	0	11
T	23	23	64	100	35	52	0	100	5
C	11	0	5	0	0	0	0	0	11

